

PCT

RAW SEQUENCE LISTING DATE: 02/22/2005
PATENT APPLICATION: US/10/501,666A TIME: 13:39:48

Input Set : A:\seq_lst_US_REV1.txt
Output Set: N:\CRF4\02222005\J501666A.raw

5 <110> APPLICANT: Stordeur, Patrick
6 Goldman, Michel
10 <120> TITLE OF INVENTION: Method to determine in vivo nucleic acid levels
14 <130> FILE REFERENCE: DECLE35.002APC
17 <140> CURRENT APPLICATION NUMBER: 10/501,666A
18 <141> CURRENT FILING DATE: 2004-07-16
21 <150> PRIOR APPLICATION NUMBER: PCT/EP03/00493
22 <151> PRIOR FILING DATE: 2003-01-20
24 <150> PRIOR APPLICATION NUMBER: EP 02447009.8
26 <151> PRIOR FILING DATE: 2002-01-18
30 <160> NUMBER OF SEQ ID NOS: 55
34 <170> SOFTWARE: PatentIn version 3.1

40 <211> LENGTH: 21
42 <212> TYPE: DNA
44 <213> ORGANISM: Artificial Sequence
48 <220> FEATURE:
50 <223> OTHER INFORMATION: Oligonucleotide

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66 <220> FEATURE:

68 <223> OTHER INFORMATION: Oligonucleotide

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38 <210> SEQ ID NO: 1

52 <400> SEQUENCE: 1

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74 <210> SEQ ID NO: 3

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88 <220> FEATURE:

90 <221> NAME/KEY: misc_feature

92 <222> LOCATION: (1)..(1)

94 <223> OTHER INFORMATION: t labelled with 6Fam

98 <220> FEATURE:

100 <221> NAME/KEY: misc_feature

102 <222> LOCATION: (25)..(25)

104 <223> OTHER INFORMATION: g labelled with Tamra-p

108 <400> SEQUENCE: 3

Dres Not Comply
Corrected Diskette Needed

(PS. 6)

21

RAW SEQUENCE LISTING DATE: 02/22/2005 TIME: 13:39:48 PATENT APPLICATION: US/10/501,666A

Input Set : A:\seq_lst_US_REV1.txt
Output Set: N:\CRF4\02222005\J501666A.raw

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118	<213> ORGANISM: Artificial Sequence	
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124	<223> OTHER INFORMATION: Oligonucleotide	
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148	<210> SEQ ID NO: 6	
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RAW SEQUENCE LISTING DATE: 02/22/2005 PATENT APPLICATION: US/10/501,666A TIME: 13:39:49

Input Set : A:\seq_lst_US_REV1.txt
Output Set: N:\CRF4\02222005\J501666A.raw

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334 <210> SEQ ID NO: 13
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Input Set : A:\seq_lst_US_REV1.txt

Output Set: N:\CRF4\02222005\J501666A.raw

336 <211> LENGTH: 20 338 <212> TYPE: DNA 340 <213> ORGANISM: Artificial Sequence 344 <220> FEATURE: 346 <223> OTHER INFORMATION: Oligonucleotide 348 <400> SEQUENCE: 13 20 349 tgaggagctg gtcaacatca 352 <210> SEQ ID NO: 14 354 <211> LENGTH: 21 356 <212> TYPE: DNA 358 <213> ORGANISM: Artificial Sequence 362 <220> FEATURE: 364 <223> OTHER INFORMATION: Oligonucleotide 366 <400> SEQUENCE: 14 21 367 caggttgatg ctccatacca t 370 <210> SEQ ID NO: 15 372 <211> LENGTH: 20 374 <212> TYPE: DNA 376 <213> ORGANISM: Artificial Sequence 380 <220> FEATURE: 382 <223> OTHER INFORMATION: Oligonucleotide 384 <220> FEATURE: 386 <221> NAME/KEY: misc feature 388 <222> LOCATION: (1)..(1) 390 <223> OTHER INFORMATION: a labelled with 6FAM 394 <220> FEATURE: 396 <221> NAME/KEY: misc feature 398 <222> LOCATION: (20)..(20) 400 <223> OTHER INFORMATION: c labelled with Tamra-p 404 <400> SEQUENCE: 15 405 aggeteege tetgeaatgg e 20 408 <210> SEQ ID NO: 16 410 <211> LENGTH: 21 412 <212> TYPE: DNA 414 <213> ORGANISM: Artificial Sequence 418 <220> FEATURE: 420 <223> OTHER INFORMATION: Oligonucleotide 422 <400> SEQUENCE: 16 423 cccagggacc tctctctaat c 21 426 <210> SEQ ID NO: 17 428 <211> LENGTH: 21 430 <212> TYPE: DNA 432 <213> ORGANISM: Artificial Sequence 436 <220> FEATURE: 438 <223> OTHER INFORMATION: Oligonucleotide 440 <400> SEQUENCE: 17 21 441 atgggctaca ggcttgtcac t 444 <210> SEQ ID NO: 18

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/501,666A

DATE: 02/22/2005
TIME: 13:39:49

Input Set : A:\seq_lst_US_REV1.txt
Output Set: N:\CRF4\02222005\J501666A.raw

448 <212> TYPE: DNA 450 <213> ORGANISM: Artificial Sequence 454 <220> FEATURE: 456 <223> OTHER INFORMATION: Oligonucleotide 458 <220> FEATURE: 460 <221> NAME/KEY: misc_feature 462 <222> LOCATION: (1)..(1) 464 <223> OTHER INFORMATION: t labelled with 6Fam 468 <220> FEATURE: 470 <221> NAME/KEY: misc feature 472 <222> LOCATION: (22)..(22) 474 <223> OTHER INFORMATION: c labelled with Tamra-p 478 <400> SEQUENCE: 18 22 479 tggcccagg cagtcagatc atc 482 <210> SEQ ID NO: 19 484 <211> LENGTH: 24 486 <212> TYPE: DNA 488 <213> ORGANISM: Artificial Sequence 492 <220> FEATURE: 494 <223> OTHER INFORMATION: Oligonucleotide 496 <400> SEQUENCE: 19 24 497 ctaattattc ggtaactgac ttga 500 <210> SEQ ID NO: 20 502 <211> LENGTH: 21 504 <212> TYPE: DNA 506 <213> ORGANISM: Artificial Sequence 510 <220> FEATURE: 512 <223> OTHER INFORMATION: Oligonucleotide 514 <400> SEQUENCE: 20 21 515 acaqttcagc catcacttgg a 518 <210> SEO ID NO: 21 520 <211> LENGTH: 24 522 <212> TYPE: DNA 524 <213> ORGANISM: Artificial Sequence 528 <220> FEATURE: 530 <223> OTHER INFORMATION: Oligonucleotide 532 <220> FEATURE: 534 <221> NAME/KEY: misc_feature 536 <222> LOCATION: (1)..(1) 538 <223> OTHER INFORMATION: t labelled with 6Fam 542 <220> FEATURE: 544 <221> NAME/KEY: misc feature 546 <222> LOCATION: (24)..(24) 548 <223> OTHER INFORMATION: c labelled with Tamra-p 552 <400> SEQUENCE: 21 24 553 tccaacgca aagcaataca tgaac 556 <210> SEQ ID NO: 22 558 <211> LENGTH: 21

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<211> 21

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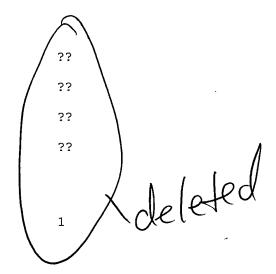
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<223> Oligonucleotide

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21



RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/501,666A

DATE: 02/22/2005 TIME: 13:39:50

Input Set : A:\seq_lst_US_REV1.txt

Output Set: N:\CRF4\02222005\J501666A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:31; N Pos. 22

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/501,666A

DATE: 02/22/2005 TIME: 13:39:50

Input Set : A:\seq_lst_US_REV1.txt

L:1292 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55

Output Set: N:\CRF4\02222005\J501666A.raw

L:803 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:1281 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:1283 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55
L:1283 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1 /
L:1285 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55
L:1285 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1 /
L:1287 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55
L:1287 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1